

PCT09

#5

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/914,001

DATE: 09/10/2001  
 TIME: 11:28:54

Input Set : A:\Methyl11.app  
 Output Set: N:\CRF3\09102001\I914001.raw

ENTERED

3 <110> APPLICANT: Wisconsin Alumni Research Foundation et al.  
 4 Pioneer Hi-Bred, International Inc.  
 5 Regents of The University of Minnesota  
 6 Kaeppler, Shawn M.  
 7 Springer, Nathan M.  
 8 Muszynski, Michael G.  
 9 Papa, Charles M.  
 11 <120> TITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Encoding Class II  
 12 DNA Methyltransferases  
 14 <130> FILE REFERENCE: WIS4987P0051PCT  
 C--> 16 <140> CURRENT APPLICATION NUMBER: US/09/914,001  
 C--> 17 <141> CURRENT FILING DATE: 2001-08-20  
 19 <160> NUMBER OF SEQ ID NOS: 90  
 21 <170> SOFTWARE: PatentIn Ver. 2.1  
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59 cagttcaagg tgcgtgtggt cctctggggg gctcttttctt ccatgtgtgct ccctaaagtat 1860
60 cctctgcccc cctatgatgt tgtagtacgt ggaggagccc ctaatgcctt ttcgcaatgt 1920
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107 cctgaggagg atacatggga accgattgat aacttgagt accgcgcga gaaattaga 1500
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145 35 40 45
147 Ala Ala Ser Ser Gly Lys Lys Pro Lys Ser Pro Pro Lys Gln Ala Lys
148 50 55 60
150 Pro Gly Lys Lys Lys Gly Asp Ala Glu Met Lys Glu Pro Val Glu Asp
151 65 70 75 80
153 Asp Val Cys Ala Glu Glu Pro Asp Glu Glu Glu Leu Ala Met Gly Glu
154 85 90 95
156 Glu Glu Ala Glu Glu Gln Ala Met Gln Glu Glu Val Val Ala Val Ala
157 100 105 110
159 Ala Gly Ser Pro Gly Lys Lys Arg Val Gly Arg Arg Asn Ala Ala Ala
160 115 120 125
162 Ala Ala Gly Asp His Glu Pro Glu Phe Ile Gly Ser Pro Val Ala Ala
163 130 135 140
165 Asp Glu Ala Arg Ser Asn Trp Pro Lys Arg Tyr Gly Arg Ser Thr Ala
166 145 150 155 160
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180 Phe Phe Arg Ala Glu Asp Thr Val Ile Asn Ser Leu Val Ser Ile Ser
181 225          230          235          240
183 Val Asp Gly His Lys His Asp Pro Arg Arg Val Phe Leu Ser Glu Glu
184          245          250          255
186 Lys Asn Asp Asn Val Leu Asp Cys Ile Ile Ser Lys Val Lys Ile Val
187          260          265          270
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192 Ser Cys Asp Leu Tyr Tyr Asp Met Ser Tyr Ser Val Ala Tyr Ser Thr
193          290          295          300
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196 305          310          315          320
198 Ser Gly Ile Ser Ser Asp Asp Val Asp Leu Glu Thr Ser Ser Ser Met
199          325          330          335
201 Pro Thr Arg Thr Ala Thr Leu Leu Asp Leu Tyr Ser Gly Cys Gly Gly
202          340          345          350
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205          355          360          365
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208          370          375          380
210 Lys Tyr Asn His Pro Gln Thr Glu Val Arg Asn Glu Lys Ala Asp Glu
211 385          390          395          400
213 Phe Leu Ala Leu Leu Lys Glu Trp Ala Val Leu Cys Lys Lys Tyr Val
214          405          410          415
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217          420          425          430
219 Asp Ser Pro Leu Asp Lys Asp Glu Phe Val Val Glu Lys Leu Val Gly
220          435          440          445
222 Ile Cys Tyr Gly Gly Ser Asp Arg Glu Asn Gly Ile Tyr Phe Lys Val
223          450          455          460
225 Gln Trp Glu Gly Tyr Gly Pro Glu Glu Asp Thr Trp Glu Pro Ile Asp
226 465          470          475          480
228 Asn Leu Ser Asp Cys Pro Gln Lys Ile Arg Glu Phe Val Gln Glu Gly
229          485          490          495
231 His Lys Arg Lys Ile Leu Pro Leu Pro Gly Asp Val Asp Val Ile Cys
232          500          505          510
234 Gly Gly Pro Pro Cys Gln Gly Ile Ser Gly Phe Asn Arg Tyr Arg Asn
235          515          520          525
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252 Trp Gly Ala Leu Ser Ser Met Val Leu Pro Lys Tyr Pro Leu Pro Thr
253      610      615      620
255 Tyr Asp Val Val Val Arg Gly Gly Ala Pro Asn Ala Phe Ser Gln Cys
256 625      630      635      640
258 Met Val Ala Tyr Asp Glu Thr Gln Lys Pro Ser Leu Lys Lys Ala Leu
259      645      650      655
261 Leu Leu Gly Asp Ala Ile Ser Asp Leu Tyr Lys Val Gln Asn His Gln
262      660      665      670
264 Pro Asn Asp Val Met Glu Tyr Gly Gly Ser Pro Lys Thr Glu Phe Gln
265      675      680      685
267 Arg Tyr Ile Arg Leu Ser Arg Lys Asp Met Leu Asp Trp Ser Phe Gly
268      690      695      700
270 Glu Gly Ala Gly Pro Asp Glu Gly Lys Leu Leu Asp His Gln Pro Leu
271 705      710      715      720
273 Arg Leu Asn Asn Asp Asp Tyr Glu Arg Val Gln Gln Ile Pro Val Lys
274      725      730      735
276 Lys Gly Ala Asn Phe Arg Asp Leu Lys Gly Val Arg Val Gly Ala Asn
277      740      745      750
279 Asn Ile Val Glu Trp Asp Pro Glu Ile Glu Arg Val Lys Leu Ser Ser
280      755      760      765
282 Gly Lys Pro Leu Val Pro Asp Tyr Ala Met Ser Phe Ile Lys Gly Lys
283      770      775      780
285 Ser Leu Lys Pro Phe Gly Arg Leu Trp Trp Asp Glu Thr Val Pro Thr
286 785      790      795      800
288 Val Val Thr Arg Ala Glu Pro His Asn Gln Val Ile Ile His Pro Thr
289      805      810      815
291 Gln Ala Arg Val Leu Thr Ile Arg Glu Asn Ala Arg Leu Gln Gly Phe
292      820      825      830
294 Pro Asp Tyr Tyr Arg Leu Phe Gly Pro Ile Lys Glu Lys Tyr Ile Gln
295      835      840      845
297 Val Gly Asn Ala Val Ala Val Pro Val Ala Arg Ala Leu Gly Tyr Cys
298      850      855      860
300 Leu Gly Gln Ala Tyr Leu Gly Glu Ser Glu Gly Ser Asp Pro Leu Tyr
301 865      870      875      880
303 Gln Leu Pro Pro Ser Phe Thr Ser Val gly Gly Arg Thr Ala Gly Gln
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VERIFICATION SUMMARY

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L:16 M:270 C: Current Application Number differs, Replaced Current Application Number

L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date